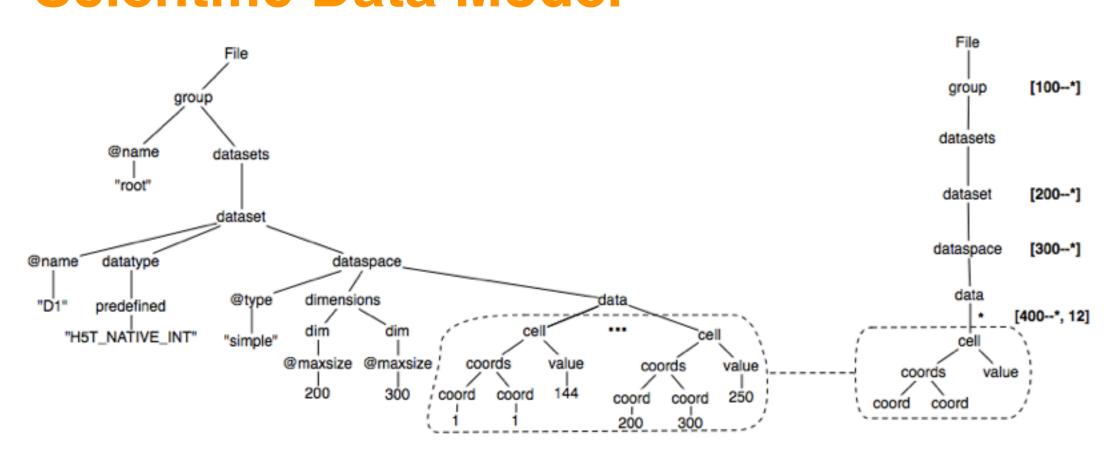


Scott Brandt (lead PI), Carlos Maltzahn (co-PI), Kleoni Ioannidou, Joe Buck, Noah Watkins UCSC – Systems Research Lab

Objectives of DAMASC:

Coalesce data management with parallel file system management to present a declarative interface to scientists for managing, querying, and analyzing extremely large data sets efficiently and predictably.

Scientific Data Model



Example representation of HDF5 file in scientific data model:
Contents represented as a node-labeled graph
Information encoded in the node labels and their nesting.

Sample schema generated by parser:Omits contents of data spacePoints to byte extents.

Data Management Services

Incremental Parsing

- Format-specific parsers expose common scientific data model
- Parse trees form structural indices into byte stream data
- Lazy, demand-driven parsing

Query Execution

- Optimized queries executed over parse trees and unmodified data
- Scientific file-formats can be modified to use DAMASC transparently from the application view-point.
- System feedback is used to tune system indices and caching behavior

Automatic Indexing

Automatic indexing of content is critical for optimized query evaluation
Content managed by DAMASC and query patterns are continuously

monitored and re-organized Provenance Tracking

- Understanding the lineage of extreme scale data is crucial as complex workflows manipulate and copy raw data
- Automatic provenance capture monitors the flow of queries in DAMASC.

File Views

- Views are expressed in the flexible, semi-structured data model used by DAMASC.
- A view is a query result, and can provide a unified view of content stored in multiple formats simultaneously.
- A view is the result of a query over the DAMASC data model
- Non-materialized views are constructed on the fly without having to create new copies of data.
- Maximum flexibility is available by allowing views to be defined on top of other views.

DAMASC

Adding Data Management Services to Parallel File Systems

Neoklis Polyzotis (co-PI), Wang-Chiew Tan (co-PI), Jeff LeFevre UCSC – Database Systems Group

Maya Gokhale (PI, LLNL), Celeste Matarazzo LLNL Jim Ahrens, John Bent, Gary Grider, James Nunez, Meghan Wingate LANL

Russ Rew

UCAR/Unidata

The POSIX IO Bottleneck

POSIX IO dominates File system interface

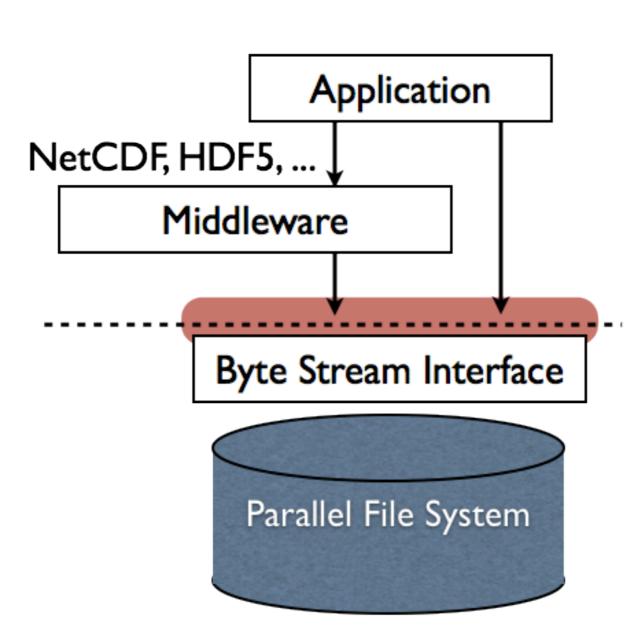
- POSIX IO does not scale
- ~50 years ago: 100MB of data at high end
- Now: up to 1 billion times more data

Performance price of POSIX IO is high

- Workload-specific interposition layers (e.g. PLFS):
- almost 1,000x speed-up

Common Workaround

• Middleware tries to make up for limitations



DAMASC Components:

Rewritten

HDF5

Tracker

Query Executor

transparently optimizes

etrieval of requested dat

Service 2:

Provenance

tracking

Parser exposes data in

structured logical model

Heavy! Do not move!

Application

Declarative

Query Executor Cache and Automatic

File Views

Byte Stream Interface

Parallel File System

Heavy! Do not move!

arser Module

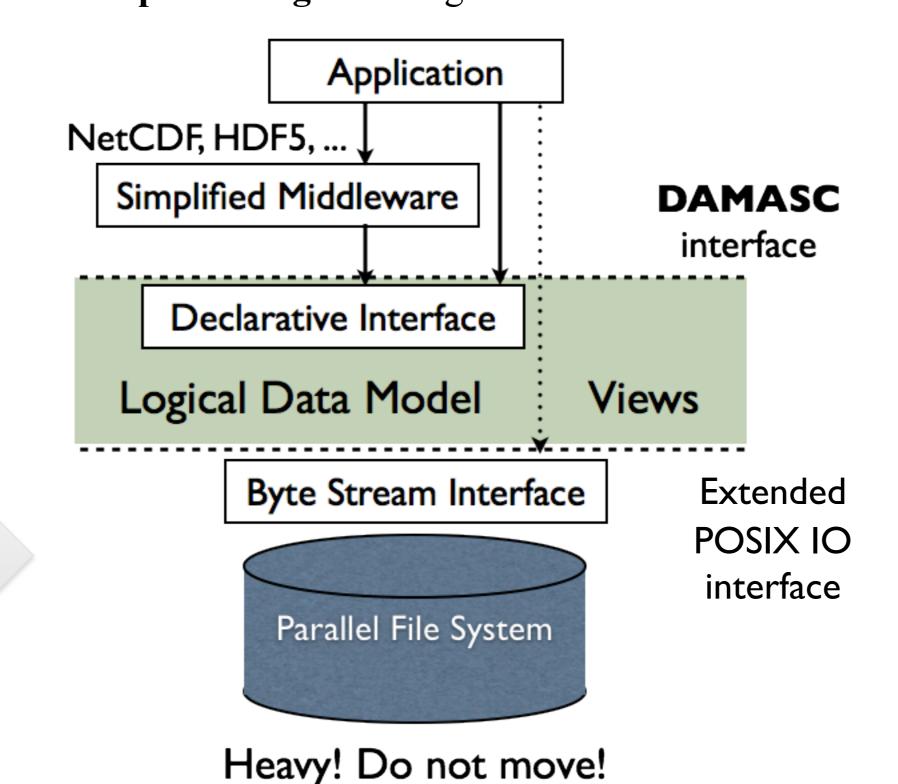
DAMASC:

DAta MAnagement in Scientific Computing

Enhance parallel file system with data services

- Declarative querying
- Views
- Automatic content indexing
- Provenance tracking

In situ processing on storage nodes



Current Work



SciHadoop:

- Intelligent input partitioning based on logical/physical mapping
- Query optimizations enabled by semantic awareness of mappers
- Use cases: HDFS and Ceph

SciZorba:

Service I:

indexing

Self-organizable

Damasc

File Views computes files

on-the-fly based on query

expressions

Extended

POSIX IO

Interface



- Declarative query processing
- Integration of NetCDF and HDF5 operators in XML queries

SciZorba

SciZorba is a system for executing scientific data queries expressed in the flexible XQuery language over data stored in the netCDF file format.

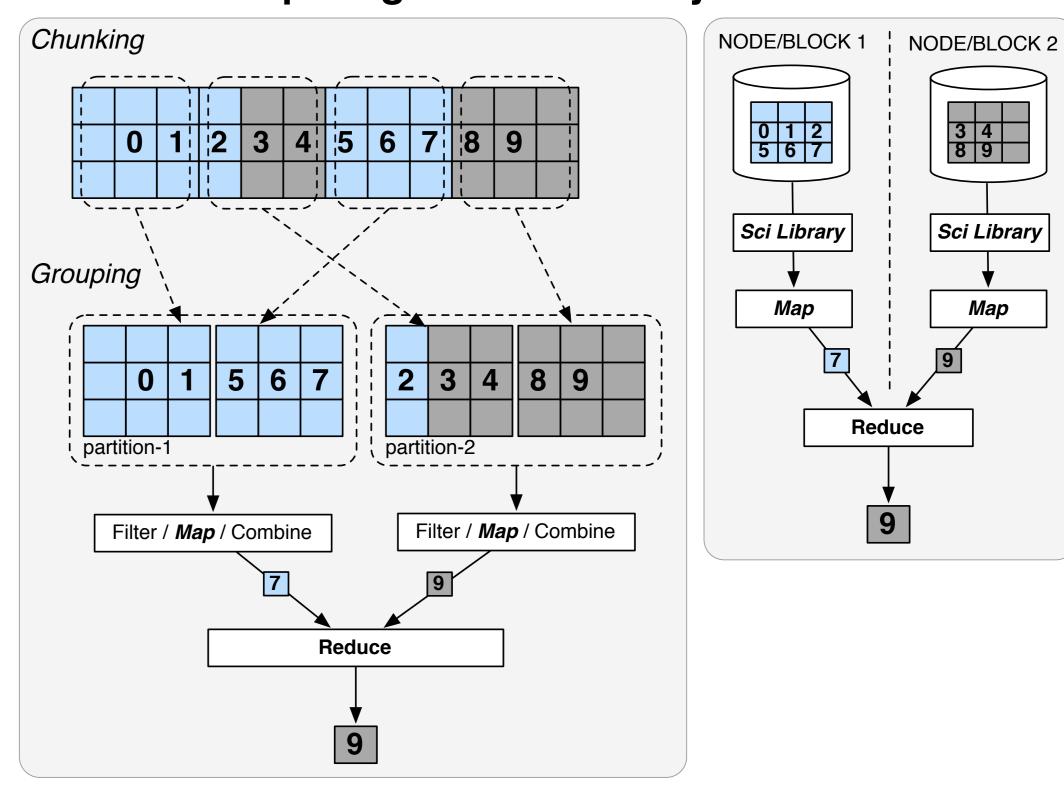
- Based on the Zorba XQuery processing engine
- Stores data in semi-structured XDM data model
- Incremental parsing loads data on-demand as required by an executing query
- Currently SciZorba runs on a single-node, and we plan to extend this to multiple cooperating nodes each running SciZorba instances that communicate to process a single query.

SciHadoop

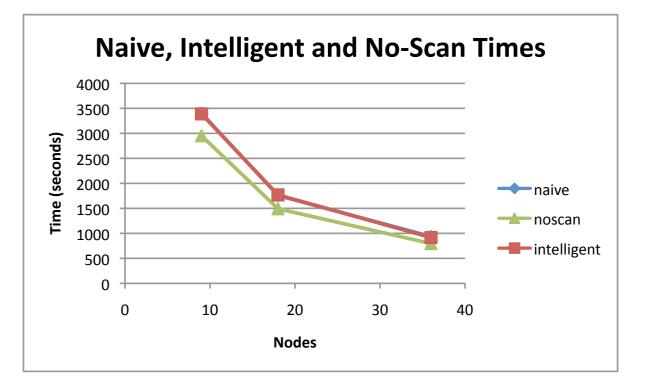
SciHadoop is a MapReduce-based framework for large-scale processing of scientific, array-based data.

- MapReduce partitions an input file's byte stream into blocks that are processed in parallel
- Reading scientific data requires passing through middleware that hides physical layout information
- These middleware libraries make it difficult to predict data locations

SciHadoop: Logical Plan vs Physical Accesses



Local Read Rate 1 0.9 0.8 0.7 0.6 0.5 0.4 0.3 0.2 0.1 0 0.0 0.2 0.1 0 0.0 0.2 0.1 0 0.0 0.2 0.1 0 0.0 0.2 0.1 0 0.0 0.2 0.1 0 0.0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.2 0.1 0 0.1 0 0.2 0 0.1 0 0.2 0 0.1 0 0



Acknowledgements

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Preliminary Results

Graphs on the left show SciHadoop enabling better locality and more efficient data access for queries over NetCDF data stored in HDFS

Future Work

- Extending support to other file formats such as HDF5 and FITS.
- Supporting write workloads by replacing HDFS with a file system such as Ceph.

Further Information

All information relating to the DAMASC project can be found at http://srl.ucsc.edu/projects/damasc